

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/757,077B
Source: TFW16
Date Processed by STIC: 2/5/07

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 02/05/2007

PATENT APPLICATION: US/10/757,077B

TIME: 14:53:27

Input Set : A:\17355CIP3SEQLIST.TXT

Output Set: N:\CRF4\02052007\J757077B.raw

4 <110> APPLICANT: Steward, Lance E.
 5 Fernandez-Salas, Ester
 6 Herrington, Todd
 7 Aoki, Kei Roger
 9 <120> TITLE OF INVENTION: Clostridial Neurotoxin Compositions and
 10 Modified Clostridial Neurotoxins
 13 <130> FILE REFERENCE: 17355CIP3 (BOT)
 15 <140> CURRENT APPLICATION NUMBER: US 10/757,077B
 16 <141> CURRENT FILING DATE: 2004-01-14
 18 <150> PRIOR APPLICATION NUMBER: US 09/910,346
 19 <151> PRIOR FILING DATE: 2001-07-20
 21 <150> PRIOR APPLICATION NUMBER: US 09/620,840
 22 <151> PRIOR FILING DATE: 2000-07-21
 24 <150> PRIOR APPLICATION NUMBER: US 10/163,106
 25 <151> PRIOR FILING DATE: 2003-06-04
 27 <160> NUMBER OF SEQ ID NOS: 83
 29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 31 <210> SEQ ID NO: 1
 32 <211> LENGTH: 7
 33 <212> TYPE: PRT
 34 <213> ORGANISM: Clostridium botulinum serotype A
 36 <400> SEQUENCE: 1
 37 Phe Glu Phe Tyr Lys Leu Leu
 38 1 5
 41 <210> SEQ ID NO: 2
 42 <211> LENGTH: 7
 43 <212> TYPE: PRT
 44 <213> ORGANISM: Rattus norvegicus
 46 <400> SEQUENCE: 2
 47 Glu Glu Lys Arg Ala Ile Leu
 48 1 5
 51 <210> SEQ ID NO: 3
 52 <211> LENGTH: 7
 53 <212> TYPE: PRT
 54 <213> ORGANISM: Rattus norvegicus
 56 <400> SEQUENCE: 3
 57 Glu Glu Lys Met Ala Ile Leu
 58 1 5
 61 <210> SEQ ID NO: 4
 62 <211> LENGTH: 7
 63 <212> TYPE: PRT
 64 <213> ORGANISM: Rattus norvegicus
 66 <400> SEQUENCE: 4

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Input Set : A:\17355CIP3SEQLIST.TXT

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67 Ser Glu Arg Asp Val Leu Leu
68 1 5
71 <210> SEQ ID NO: 5
72 <211> LENGTH: 7
73 <212> TYPE: PRT
74 <213> ORGANISM: Rattus norvegicus
76 <400> SEQUENCE: 5
77 Val Asp Thr Gln Val Leu Leu
78 1 5
81 <210> SEQ ID NO: 6
82 <211> LENGTH: 7
83 <212> TYPE: PRT
84 <213> ORGANISM: Mus musculus
86 <400> SEQUENCE: 6
87 Ala Glu Val Gln Ala Leu Leu
88 1 5
91 <210> SEQ ID NO: 7
92 <211> LENGTH: 7
93 <212> TYPE: PRT
94 <213> ORGANISM: Xenopus laevis
96 <400> SEQUENCE: 7
97 Ser Asp Lys Gln Asn Leu Leu
98 1 5
101 <210> SEQ ID NO: 8
102 <211> LENGTH: 7
103 <212> TYPE: PRT
104 <213> ORGANISM: Gallus gallus
106 <400> SEQUENCE: 8
107 Ser Asp Arg Gln Asn Leu Ile
108 1 5
111 <210> SEQ ID NO: 9
112 <211> LENGTH: 7
113 <212> TYPE: PRT
114 <213> ORGANISM: Ovis aries
116 <400> SEQUENCE: 9
117 Ala Asp Thr Gln Val Leu Met
118 1 5
121 <210> SEQ ID NO: 10
122 <211> LENGTH: 7
123 <212> TYPE: PRT
124 <213> ORGANISM: Homo sapiens
126 <400> SEQUENCE: 10
127 Ser Asp Lys Asn Thr Leu Leu
128 1 5
131 <210> SEQ ID NO: 11
132 <211> LENGTH: 7
133 <212> TYPE: PRT
134 <213> ORGANISM: Homo sapiens
136 <400> SEQUENCE: 11

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Input Set : A:\17355CIP3SEQLIST.TXT

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137 Ser Gln Ile Lys Arg Leu Leu

138 1 5

141 <210> SEQ ID NO: 12

142 <211> LENGTH: 7

143 <212> TYPE: PRT

144 <213> ORGANISM: Homo sapiens

146 <400> SEQUENCE: 12

147 Ala Asp Thr Gln Ala Leu Leu

148 1 5

151 <210> SEQ ID NO: 13

152 <211> LENGTH: 7

153 <212> TYPE: PRT

154 <213> ORGANISM: Saccharomyces cerevisiae

156 <400> SEQUENCE: 13

157 Asn Glu Gln Ser Pro Leu Leu

158 1 5

161 <210> SEQ ID NO: 14

162 <211> LENGTH: 12

163 <212> TYPE: PRT

164 <213> ORGANISM: Clostridium botulinum serotype A

166 <400> SEQUENCE: 14

167 Met Pro Phe Val Asn Lys Gln Phe Asn Tyr Lys Asp

168 1 5 10

171 <210> SEQ ID NO: 15

172 <211> LENGTH: 11

173 <212> TYPE: PRT

174 <213> ORGANISM: Clostridium botulinum serotype A

176 <400> SEQUENCE: 15

177 Pro Phe Val Asn Lys Gln Phe Asn Tyr Lys Asp

178 1 5 10

181 <210> SEQ ID NO: 16

182 <211> LENGTH: 4

183 <212> TYPE: PRT

184 <213> ORGANISM: Clostridium botulinum serotype A

186 <400> SEQUENCE: 16

187 Met Tyr Lys Asp

188 1

191 <210> SEQ ID NO: 17

192 <211> LENGTH: 7

193 <212> TYPE: PRT

194 <213> ORGANISM: Artificial Sequence

196 <220> FEATURE:

197 <221> NAME/KEY: SITE

198 <222> LOCATION: (1)...(7)

199 <223> OTHER INFORMATION: Consensus sequence for Leucine-based motif.

201 <221> NAME/KEY: VARIANT

202 <222> LOCATION: (1)...(1)

203 <223> OTHER INFORMATION: Xaa is any amino acid.

W--> 205 <221> VARIANT

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Input Set : A:\17355CIP3SEQLIST.TXT

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206 <222> LOCATION: (3)...(5)
207 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 209 <400> 17
W--> 210 Xaa Asp Xaa Xaa Xaa Leu Leu
211 1 5
214 <210> SEQ ID NO: 18
215 <211> LENGTH: 7
216 <212> TYPE: PRT
217 <213> ORGANISM: Artificial Sequence
219 <220> FEATURE:
220 <221> NAME/KEY: SITE
221 <222> LOCATION: (1)...(7)
222 <223> OTHER INFORMATION: Consensus sequence for Leucine-based motif.
224 <221> NAME/KEY: VARIANT
225 <222> LOCATION: (1)...(1)
226 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 228 <221> VARIANT
229 <222> LOCATION: (3)...(5)
230 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 232 <400> 18
W--> 233 Xaa Glu Xaa Xaa Xaa Leu Leu
234 1 5
237 <210> SEQ ID NO: 19
238 <211> LENGTH: 7
239 <212> TYPE: PRT
240 <213> ORGANISM: Artificial Sequence
242 <220> FEATURE:
243 <221> NAME/KEY: SITE
244 <222> LOCATION: (1)...(7)
245 <223> OTHER INFORMATION: Consensus sequence for Leucine-based motif.
247 <221> NAME/KEY: VARIANT
248 <222> LOCATION: (1)...(1)
249 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 251 <221> VARIANT
252 <222> LOCATION: (3)...(5)
253 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 255 <400> 19
W--> 256 Xaa Asp Xaa Xaa Xaa Leu Ile
257 1 5
260 <210> SEQ ID NO: 20
261 <211> LENGTH: 7
262 <212> TYPE: PRT
263 <213> ORGANISM: Artificial Sequence
265 <220> FEATURE:
266 <221> NAME/KEY: SITE
267 <222> LOCATION: (1)...(7)
268 <223> OTHER INFORMATION: Consensus sequence for Leucine-based motif.
270 <221> NAME/KEY: VARIANT
271 <222> LOCATION: (1)...(1)

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272 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 274 <221> VARIANT
275 <222> LOCATION: (3)...(5)
276 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 278 <400> 20
W--> 279 Xaa Asp Xaa Xaa Xaa Leu Met
280 1 5
283 <210> SEQ ID NO: 21
284 <211> LENGTH: 7
285 <212> TYPE: PRT
286 <213> ORGANISM: Artificial Sequence
288 <220> FEATURE:
289 <221> NAME/KEY: SITE
290 <222> LOCATION: (1)...(7)
291 <223> OTHER INFORMATION: Consensus sequence for Leucine-based motif.
293 <221> NAME/KEY: VARIANT
294 <222> LOCATION: (1)...(1)
295 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 297 <221> VARIANT
298 <222> LOCATION: (3)...(5)
299 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 301 <400> 21
W--> 302 Xaa Glu Xaa Xaa Xaa Leu Ile
303 1 5
306 <210> SEQ ID NO: 22
307 <211> LENGTH: 7
308 <212> TYPE: PRT
309 <213> ORGANISM: Artificial Sequence
311 <220> FEATURE:
312 <221> NAME/KEY: SITE
313 <222> LOCATION: (1)...(7)
314 <223> OTHER INFORMATION: Consensus sequence for Leucine-based motif.
316 <221> NAME/KEY: VARIANT
317 <222> LOCATION: (1)...(1)
318 <223> OTHER INFORMATION: Xaa is any amino acid..
W--> 320 <221> VARIANT
321 <222> LOCATION: (3)...(5)
322 <223> OTHER INFORMATION: Xaa is any amino acid.
W--> 324 <400> 22
W--> 325 Xaa Glu Xaa Xaa Xaa Ile Leu
326 1 5
329 <210> SEQ ID NO: 23
330 <211> LENGTH: 7
331 <212> TYPE: PRT
332 <213> ORGANISM: Artificial Sequence
334 <220> FEATURE:
335 <221> NAME/KEY: SITE
336 <222> LOCATION: (1)...(7)
337 <223> OTHER INFORMATION: Consensus sequence for Leucine-based motif.

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/757,077B

DATE: 02/05/2007
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Input Set : A:\17355CIP3SEQLIST.TXT
Output Set: N:\CRF4\02052007\J757077B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:17; Xaa Pos. 1,3,4,5 ✓
Seq#:18; Xaa Pos. 1,3,4,5 ✓
Seq#:19; Xaa Pos. 1,3,4,5 ✓
Seq#:20; Xaa Pos. 1,3,4,5 ✓
Seq#:21; Xaa Pos. 1,3,4,5 ✓
Seq#:22; Xaa Pos. 1,3,4,5 ✓
Seq#:23; Xaa Pos. 1,3,4,5 ✓
Seq#:24; Xaa Pos. 2,3,4

VERIFICATION SUMMARY

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Input Set : A:\17355CIP3SEQLIST.TXT

Output Set: N:\CRF4\02052007\J757077B.raw

L:205 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:209 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:228 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:232 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:251 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:255 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:274 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:278 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:297 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21
L:301 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21
L:302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:320 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:324 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:325 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:343 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:347 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:348 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:366 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:370 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:905 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:53
L:926 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:54
L:951 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:55
L:972 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:56
L:976 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:56
L:1001 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:57
L:1022 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:58
L:1047 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:59
L:1068 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:60
L:1093 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:61
L:1114 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:62
L:1139 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:63
L:1143 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:63
L:1164 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:64
L:1189 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:65
L:1210 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:66
L:1235 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:67
L:1256 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:68
L:1281 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:69
L:1285 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:69
L:1306 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:70
L:1331 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:71
L:1352 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:72
L:1377 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:73

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L:1398 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:74
L:1402 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:74
L:1427 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:75
L:1448 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:76
L:1473 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:77
L:1494 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:78
L:1519 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:79
L:1540 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:80